



Figure S3 Gene expression levels in control and RNAi knockdown measured by quantitative RT-PCR. Ubiquitous expression of RNAi against a subset of (A) positive candidates and (B) negative candidates result in varying degrees of knockdown. The mean transcript levels were calculated from three biological replicates using $\Delta\Delta C_T$ method and normalized to *rp49* mRNA levels. Controls are RNAi to GFP. Error bars represent standard deviation.